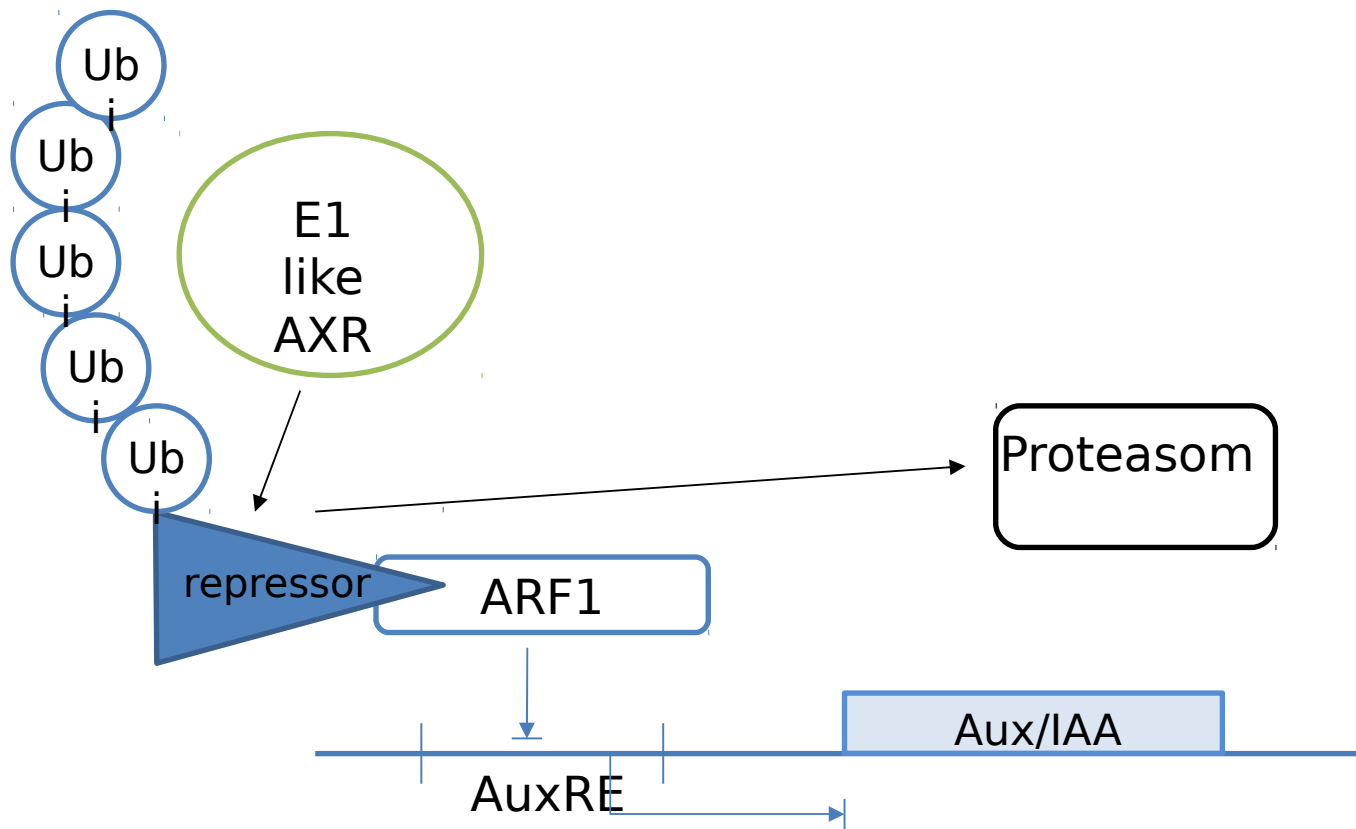


The TIR1 protein of Arabidopsis functions in auxin response and is related to human SKP2 and yeast Grr1p

Max Ruegger, Elizabeth Dewey, William M. Gray, Lawrence
Hobbie, Jocelin Turner and Mark Estelle



Ziel:

Identification of a new gene
called TRANSPORT INHIBITOR
RESPONSE 1 (TIR1)

Table 1. Segregation of CPD resistance in a *tir1-2* × wild-type F_2 population

F ₂ root length (mm)	F ₂ genotype ¹		
	<i>tir1/tir1</i> ²	<i>tir1/+</i> ²	<i>+/+</i> ²
17	2		
16	5		
15	1		
14			
13			
12		1	
11			
10		6	
9		6	
8		7	6
7		3	4
6		2	4
5			
4		1	
3		1	
Totals	8	27	14

¹Genotype determined by analyzing CPD resistance in F₃ plants.

²Number of seedlings.

Vergleich der Wurzellängen von homo- und heterozygoten Pflanzen auf CPD-Medium

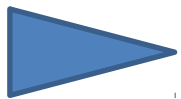


Mutation wirkt semidominant

Table 2. *CPD resistance in tir homozygous and heterozygous plants*

Line	Root length (mm)	±S.E.	No.
+/+	7.8	0.1	32
<i>tir1-1/tir1-1</i>	15.7	0.3	33
<i>tir1-3/tir1-3</i>	15.8	0.4	34
<i>tir1-7/tir1-7</i>	13.0	0.5	33
<i>tir1-1/+</i>	10.9	0.3	15
<i>tir1-3/+</i>	12.5	0.3	17
<i>tir1-7/+</i>	12.6	0.2	17

Vergleich von Mutationen von unterschiedlichen Allelen



Bei allen Allelen ist die Semidominanz nachweisbar

Besteht die Resistenz auf Grund
eines veränderten
Transportwegs, oder tritt generell
keine Auxinantwort auf??

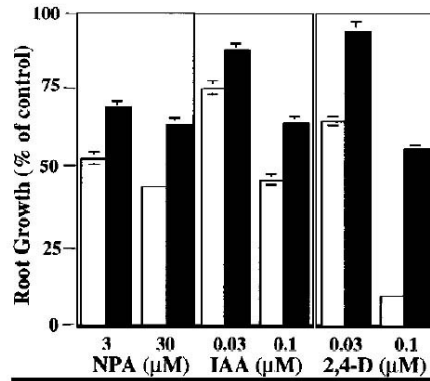
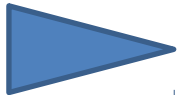


Figure 1. The roots of wild-type (open bars) and *tir1-1* (solid bars) seedlings are resistant to the growth-inhibiting properties of NPA, IAA, and 2,4-D. Seeds were germinated on nutrient medium. After 3 days, seedlings were transferred to media containing the indicated compound. Five days later, new root growth was measured and plotted as a percentage of root growth on medium without compound. Bars represent standard errors. Absence of bar indicates error less than thickness of line ($n = 12$).

Vergleich des Wurzelwachstums
von Wildtyp mit Mutante



Signifikante Unterschiede
erkennbar



Beim Mutant liegt ein genereller Defekt in der
Auxinantwort vor

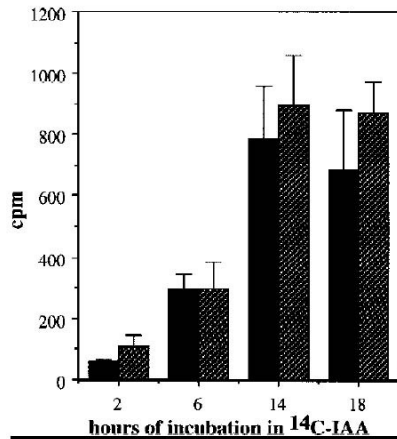
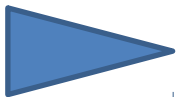


Figure 2. Polar auxin transport is similar in wild-type (solid bars) and *tir1-1* (hatched bars) inflorescence stems. Two and one-half centimeters of stem was excised, and the apical end was placed in a nutrient solution containing $1 \mu\text{M}$ [^{14}C]IAA for the times indicated. The amount of radioactive IAA transported to the basal end of the stem was assayed by liquid scintillation. Each column represents the mean of three replicates; the bar represents the standard error of the mean.

Vergleich des polaren
Auxintransport von Wildtyp und
Mutante

▶ Keine signifikanten Unterschiede
erkennbar



Es liegt ein genereller Defekt in der Auxinantwort vor

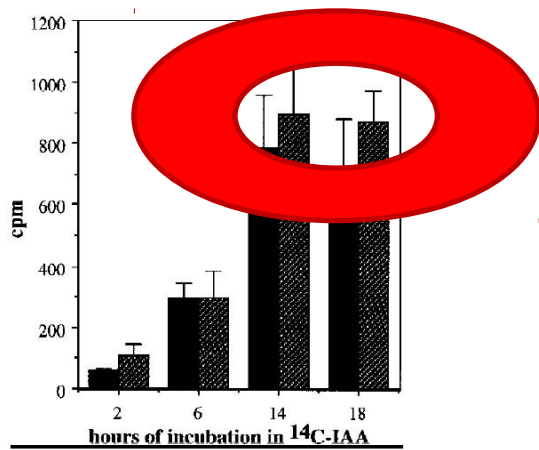


Figure 2. Polar auxin transport is similar in wild-type (solid bars) and *tir1-1* (hatched bars) inflorescence stems. Two and one-half centimeters of stem was excised, and the apical end was placed in a nutrient solution containing $1 \mu\text{M}$ [^{14}C]IAA for the times indicated. The amount of radioactive IAA transported to the basal end of the stem was assayed by liquid scintillation. Each column represents the mean of three replicates; the bar represents the standard error of the mean.

Vergleich des polaren Auxintransport von Wildtyp und Mutante



Keine signifikanten Unterschiede erkennbar



Es liegt ein genereller Defekt in der Auxinantwort vor

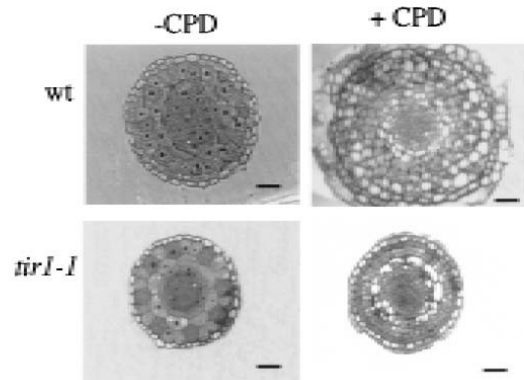


Figure 3. Cell proliferation in response to CPD is reduced in the roots of *tir1-1* seedlings. Seedlings were grown for 7 days on nutrient medium plus or minus 5 μ M CPD, fixed, and embedded in Spurr's (Ruegger et al. 1997). Root cross sections 300 μ m from the root tip are shown. Bars, 20 μ m.

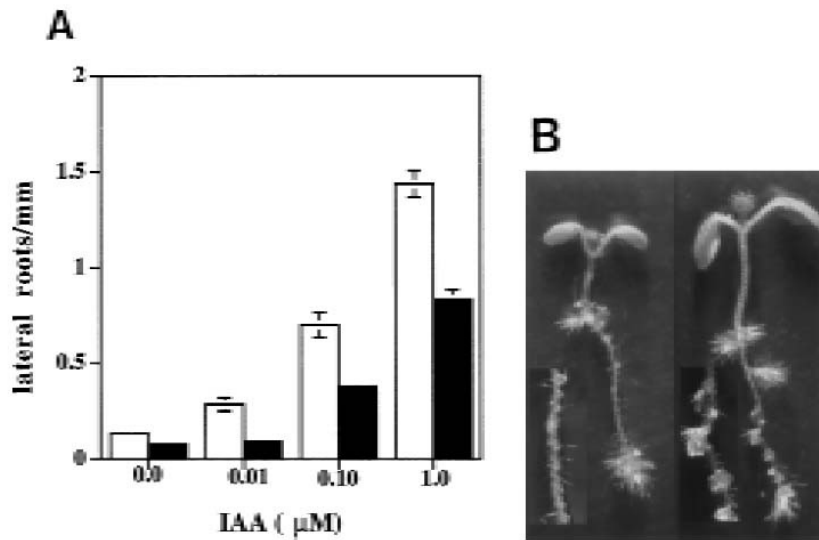
Vergleich von Zellteilungsrate in der Wurzel

▶ Bei unbehandelten Pflanzen kein Unterschied erkennbar

▶ Bei behandelten Pflanzen weißt Wildtyp vermehrte Zellzahlen auf



tir1-1 Mutante reagiert nicht auf Auxin als Zellteilungssignal



Vergleich von
Seitenwurzelbildung bei
Mutante und Wildtyp unter IAA-
Einfluss

▶ Wildtyp bildet mehr
Seitenwurzeln

Figure 4. The *tir1-1* mutant is deficient in IAA-induction of lateral roots. (A) Eight-millimeter root segments were excised from 5-day-old wild-type (open bar) and *tir1-1* (solid bar) seedlings grown on nutrient medium and transferred to medium with IAA. Lateral roots were counted after 5 days by use of a dissecting microscope. Bars represent standard errors. Absence of bar indicates error less than thickness of line ($n = 10$). (B) Ten-day-old wild-type (left) and *tir1-1* (right) seedlings grown on $0.5 \mu\text{M}$ 2,4-D. (Insets) Higher magnification images of roots.

▶ Mutante reagiert nicht auf Auxin als Zellteilungs- und Zelldifferenzierungssignal

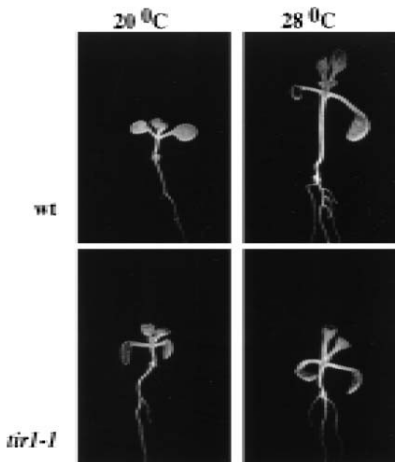
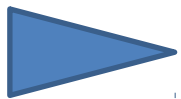


Figure 5. Hypocotyl elongation in response to elevated temperature is reduced in *tir1-1* seedlings compared with wild-type seedlings. Seedlings were grown on vertically oriented agar medium for 9 days at the two temperatures.

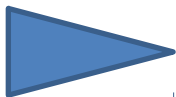
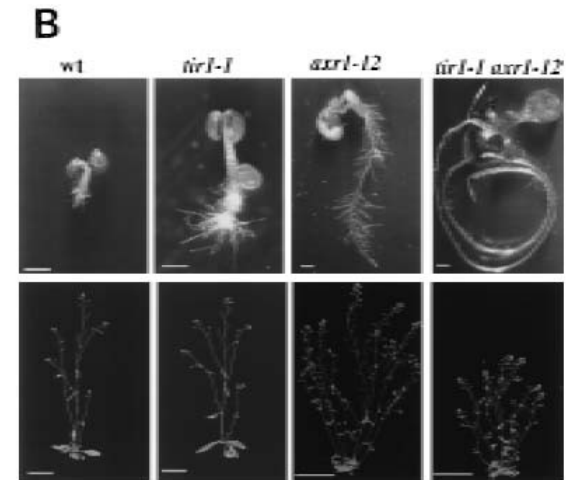
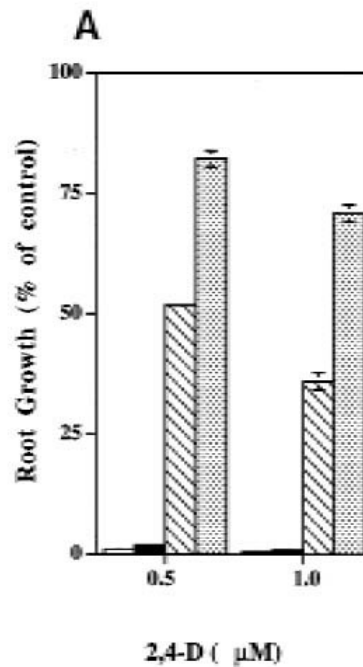
Vergleich von Hypocotyl-Elongation zwischen Wildtyp und Mutante

▶ Wildtyp zeigt mehr Hypocotyl-Elongation



Mutante reagiert nicht auf Auxin als Zellstreckungssignal

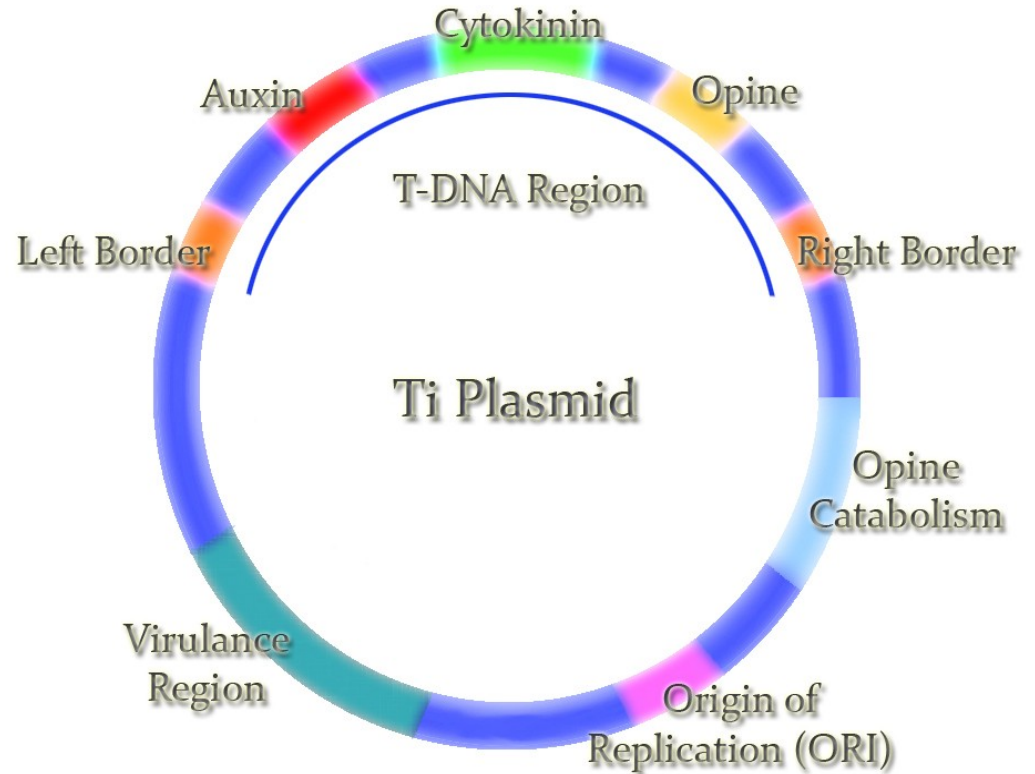
Figure 6. *tir1-1* and *axr1-12* have synergistic effects on auxin response and plant morphology. (A) The *tir1-1* mutation decreased auxin response in an *axr1-12* background at a concentration at which *tir1-1* has no effect by itself. Seedlings were grown and treated as described for Fig. 1. (Open bars) Wild type; (solid bars) *tir1*; (hatched bars) *axr1*; (stippled bars) *axr1 tir1*. Bars represent standard error and absence of bar indicates error less than thickness of line. (B) Seedlings (*top row*) were photographed after 12 days of growth on 1 μ M 2,4-D. Bar, 2.5 mm. Mature plants (*bottom row*) were photographed 35 days after germination in soil. Bar, 5 cm.



TIR1 ist am AXR1 vermittelten Prozess beteiligt

T-DNA

- stammen aus Agrobacterium
- führen zum knockout des Genes
- mittlerweile alle Insertionsstellen bekannt



Quelle: wikipedia.org

A

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TIR1      9  DEEVEHEHESFIQLDKERNLSVCHSYELCAWCRKKF
SKP2     112  SPDOHSLGHSICLCL.PEHLKPSGAEHRGRCA..SDESEN
Grr1     320  LRSSEIHELDKLNQKYGVKPLTSEILAMIEV.KIL.YY
C02F5.7   60  LKKEVLKPSFSLDT.KALCRGAQVCHSSIL..LDGSNN
UPO       50  LPPILDRVIAFLPP.PAFRTRCVHRFTSCL..FSNTCL
FIM       47  LKOKHIDRLIATLPP.PAFRRRSKVRRAVSE..FSTTPE
Cyclin F  35  LQELVDFHLKWLSEV.LALAPRAHSQKDDND.NHASTW
Cdc4     278  LKPSISLALNYLQFE.LIINLGSQSNKLR.KSTSEW

Consensus  LP-ELL--IP--L----Da--a--VCK-WY-La-----a-
              V  VL              R  I
              T

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B

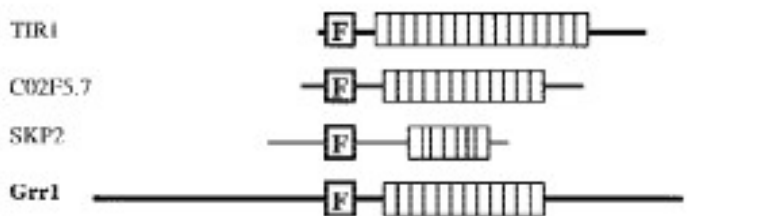
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95  WIRNHSSTYTWRFFIRK
114  RM.VVITDDLELIARSKNFKVLVLSSCEGFSTD
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202  ALERENTREPNIKSEHNRRAVPLEKIA
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363  GLVSVSMCCPKDESUYECROMTNA
388  ALITIANRNPNTRELCIIEPKAPDYLTLEPLDI
423  GFGAWEHCKDRRSLSGLLTDKVFTE
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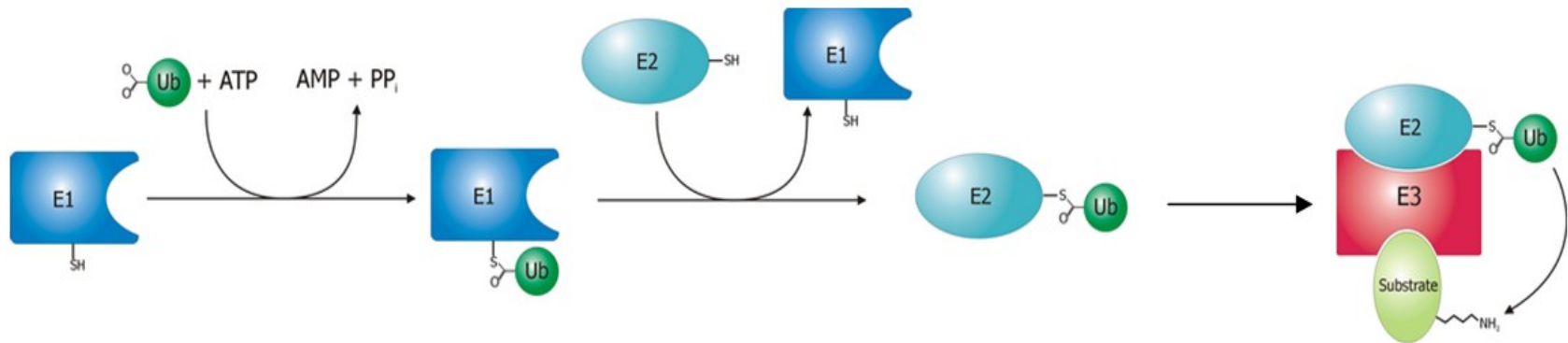
Con1 :L..a...C..L..a.....

C



- F-Box und LRR (LEUCIN-RICH-REPEATS)
- Ähnlichkeiten zu Grrp1 (Hefe), SKP2 (Mensch)
- F-Box und LRR bilden Protein-Protein-Interaktionen

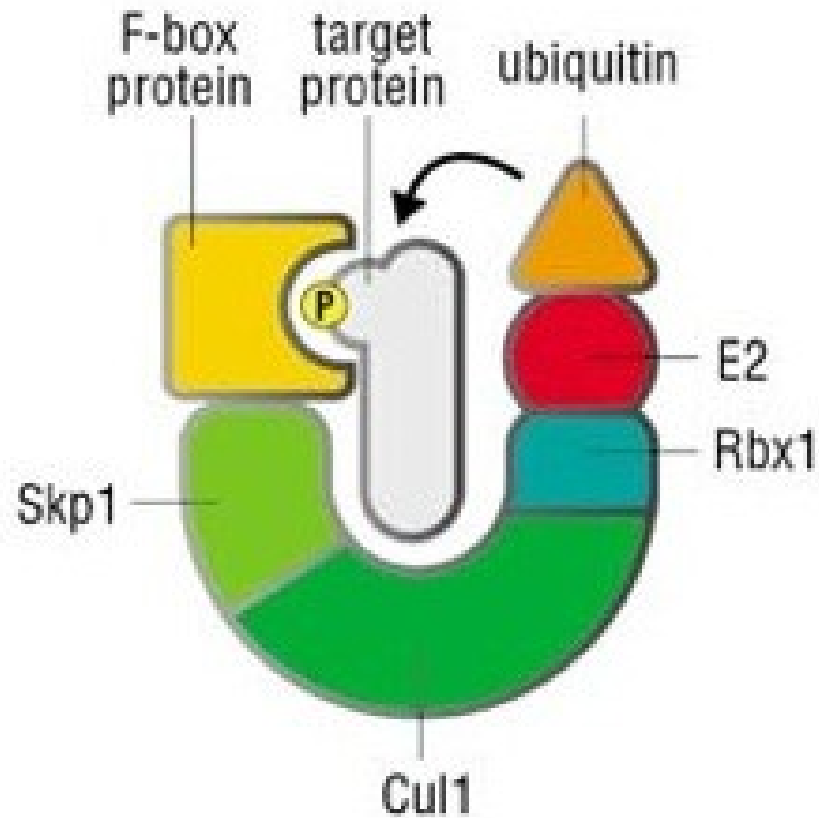
Übersicht Ubiquitinierung



- schon bekannt beim Menschen

SCF Komplex

- Skp, Cullin, F-box containing complex



Hypothese

TIR1 ist Teil der Ubiquitin-protein Ligase (E3) und bewirkt Transfer von Ubiquitin auf Zielprotein.

